

#7  
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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/462,671

DATE: 06/07/2001  
TIME: 11:26:48

Input Set : A:\PTO.txt  
Output Set: C:\CRF3\06072001\I462671.raw

ENTERED

3 <110> APPLICANT: YLIHONKO, Kristiina  
4 HAKALA, Juha  
5 KUNNARI, Tero  
7 <120> TITLE OF INVENTION: HYBRID ANTHRACYCLINES FROM GENETICALLY ENGINEERED  
STREPTOMYCES GALILAEUS  
8 STRAINS  
10 <130> FILE REFERENCE: 1574/48472  
12 <140> CURRENT APPLICATION NUMBER: 09/462,671  
13 <141> CURRENT FILING DATE: 2000-01-11  
15 <160> NUMBER OF SEQ ID NOS: 2  
17 <170> SOFTWARE: PatentIn version 3.0  
19 <210> SEQ ID NO: 1  
20 <211> LENGTH: 23  
21 <212> TYPE: DNA  
22 <213> ORGANISM: Streptomyces lividans  
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25 acatgtccga acgcacatcgccg 23  
28 <210> SEQ ID NO: 2  
29 <211> LENGTH: 23  
30 <212> TYPE: DNA  
31 <213> ORGANISM: Streptomyces lividans  
33 <400> SEQUENCE: 2  
34 agcagcgggc gggagagacg atg 23

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/462,671

DATE: 06/07/2001

TIME: 11:26:49

Input Set : A:\PTO.txt

Output Set: C:\CRF3\06072001\I462671.raw

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7/7/01

RAW SEQUENCE LISTING DATE: 06/07/2001  
PATENT APPLICATION: US/09/685,010 TIME: 11:23:29

Input Set : A:\401c1.app.txt  
Output Set: C:\CRF3\06072001\I685010.raw

P.S

4 <110> APPLICANT: Turley, Eva A.  
 5 Cruz, Tony F.  
 7 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING  
 8 CELLULAR RESPONSE TO INJURY AND OTHER PROLIFERATING CELL  
 9 DISORDERS REGULATED BY HYALADHERIN AND HYALURONANS  
 12 <130> FILE REFERENCE: 910130.401C1  
 14 <140> CURRENT APPLICATION NUMBER: US 09/685,010  
**C--> 15 <141> CURRENT FILING DATE: 2001-05-23**  
 17 <160> NUMBER OF SEQ ID NOS: 72  
 19 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
 21 <210> SEQ ID NO: 1  
 22 <211> LENGTH: 8  
 23 <212> TYPE: PRT  
 24 <213> ORGANISM: Artificial Sequence  
 26 <220> FEATURE:  
 27 <223> OTHER INFORMATION: Peptide that binds a hyalauronan  
 29 <221> NAME/KEY: VARIANT  
 30 <222> LOCATION: (1)...(5)  
 31 <223> OTHER INFORMATION: Xaa = any amino acid  
 33 <221> NAME/KEY: HELIX  
 34 <222> LOCATION: (1)...(5)  
 35 <223> OTHER INFORMATION: Alpha-helix  
 37 <221> NAME/KEY: VARIANT  
 38 <222> LOCATION: (6)...(8)  
 39 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
 41 <400> SEQUENCE: 1  
**W-1> 42 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa**  
 43 1 5  
 45 <210> SEQ ID NO: 2  
 46 <211> LENGTH: 11  
 47 <212> TYPE: PRT  
 48 <213> ORGANISM: Artificial Sequence  
 50 <220> FEATURE:  
 51 <223> OTHER INFORMATION: Peptide that binds a hyalauronan  
 53 <221> NAME/KEY: VARIANT  
 54 <222> LOCATION: (1)...(5)  
 55 <223> OTHER INFORMATION: Xaa = any amino acid  
 57 <221> NAME/KEY: HELIX  
 58 <222> LOCATION: (1)...(5)  
 59 <223> OTHER INFORMATION: Alpha-helix  
 61 <221> NAME/KEY: VARIANT  
 62 <222> LOCATION: (6)...(6)  
 63 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
 65 <221> NAME/KEY: VARIANT  
 66 <222> LOCATION: (7)...(7)  
 67 <223> OTHER INFORMATION: Xaa = Hydrophobic or neutral amino acid consisting  
 68 of I,L,V,Q,S

ENTERED

Input Set : A:\401c1.app.txt  
Output Set: C:\CRF3\06072001\I685010.raw

70 <221> NAME/KEY: VARIANT  
71 <222> LOCATION: (8)...(9)  
72 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
74 <221> NAME/KEY: VARIANT  
75 <222> LOCATION: (10)...(10)  
76 <223> OTHER INFORMATION: Xaa = Hydrophobic or neutral amino acid consisting  
77 of I,L,V,Q,S  
79 <221> NAME/KEY: VARIANT  
80 <222> LOCATION: (11)...(11)  
81 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
83 <400> SEQUENCE: 2/  
84 Xaa  
85 1 5 10  
87 <210> SEQ ID NO: 3  
88 <211> LENGTH: 12  
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90 <213> ORGANISM: Artificial Sequence  
92 <220> FEATURE:  
93 <223> OTHER INFORMATION: Peptide that binds a hyaluronan  
95 <221> NAME/KEY: VARIANT  
96 <222> LOCATION: (1)...(5)  
97 <223> OTHER INFORMATION: Xaa = any amino acid  
99 <221> NAME/KEY: HELIX  
100 <222> LOCATION: (1)...(5)  
101 <223> OTHER INFORMATION: Alpha-helix  
103 <221> NAME/KEY: VARIANT  
104 <222> LOCATION: (6)...(6)  
105 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
107 <221> NAME/KEY: VARIANT  
108 <222> LOCATION: (7)...(7)  
109 <223> OTHER INFORMATION: Xaa = Hydrophobic or neutral amino acid consisting  
110 of I,L,V,Q,S  
112 <221> NAME/KEY: VARIANT  
113 <222> LOCATION: (8)...(8) / /  
114 <223> OTHER INFORMATION: Xaa = Lysinse or Arginine  
116 <221> NAME/KEY: VARIANT  
117 <222> LOCATION: (9)...(9) /  
118 <223> OTHER INFORMATION: Xaa = Hydrophobic or neutral amino acid consisting  
119 of I,L,V,Q,S  
121 <221> NAME/KEY: VARIANT  
122 <222> LOCATION: (10)...(12)  
123 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
125 <400> SEQUENCE: 3 / / /  
126 Xaa  
127 1 5 10  
129 <210> SEQ ID NO: 4  
130 <211> LENGTH: 10  
131 <212> TYPE: PRT  
132 <213> ORGANISM: Artificial Sequence

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Input Set : A:\401c1.app.txt  
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134 <220> FEATURE:  
135 <223> OTHER INFORMATION: Peptide that binds a hyaluronan  
137 <221> NAME/KEY: VARIANT  
138 <222> LOCATION: (1)...(5)  
139 <223> OTHER INFORMATION: Xaa = any amino acid  
141 <221> NAME/KEY: HELIX  
142 <222> LOCATION: (1)...(5)  
143 <223> OTHER INFORMATION: Alpha-helix  
145 <221> NAME/KEY: VARIANT  
146 <222> LOCATION: (6)...(6)  
147 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
149 <221> NAME/KEY: VARIANT  
150 <222> LOCATION: (7)...(7)  
151 <223> OTHER INFORMATION: Xaa = Hydrophobic or neutral amino acid consisting  
152 of I,L,V,Q,S  
154 <221> NAME/KEY: VARIANT  
155 <222> LOCATION: (8)...(10)  
156 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
158 <400> SEQUENCE:- 4' / /  
W-> 159 Xaa  
160 1 5 10  
162 <210> SEQ ID NO: 5  
163 <211> LENGTH: 9  
164 <212> TYPE: PRT  
165 <213> ORGANISM: Artificial Sequence  
167 <220> FEATURE:  
168 <223> OTHER INFORMATION: Peptide that binds a hyaluronan  
170 <221> NAME/KEY: VARIANT  
171 <222> LOCATION: (1)...(5)  
172 <223> OTHER INFORMATION: Xaa = Any amino acid  
174 <221> NAME/KEY: HELIX  
175 <222> LOCATION: (1)...(5)  
176 <223> OTHER INFORMATION: Alpha-helix  
178 <221> NAME/KEY: VARIANT  
179 <222> LOCATION: (6)...(6)  
180 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
182 <221> NAME/KEY: VARIANT  
183 <222> LOCATION: (7)...(7)  
184 <223> OTHER INFORMATION: Xaa = Hydrophobic or neutral amino acid consisting  
185 of I,L,V,Q,S  
187 <221> NAME/KEY: VARIANT  
188 <222> LOCATION: (8)...(9)  
189 <223> OTHER INFORMATION: Xaa = Lysine or Arginine  
191 <400> SEQUENCE: 5'  
W-> 192 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa  
193 1 5  
195 <210> SEQ ID NO: 6  
196 <211> LENGTH: 7  
197 <212> TYPE: PRT

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/685,010

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TIME: 11:23:29

Input Set : A:\401c1.app.txt  
Output Set: C:\CRF3\06072001\I685010.raw

198 <213> ORGANISM: Artificial Sequence  
200 <220> FEATURE:  
201 <223> OTHER INFORMATION: Peptide that binds a hyaluronan  
203 <221> NAME/KEY: HELIX  
204 <222> LOCATION: (1)...(5)  
205 <223> OTHER INFORMATION: Alpha-helix  
207 <400> SEQUENCE: 6  
208 Met Met Thr Val Leu Lys Arg  
209 1 5  
211 <210> SEQ ID NO: 7  
212 <211> LENGTH: 11  
213 <212> TYPE: PRT  
214 <213> ORGANISM: Artificial Sequence  
216 <220> FEATURE:  
217 <223> OTHER INFORMATION: Peptide that binds a hyaluronan  
219 <221> NAME/KEY: HELIX  
220 <222> LOCATION: (1)...(5)  
221 <223> OTHER INFORMATION: Alpha-helix  
223 <400> SEQUENCE: 7  
224 Met Met Thr Val Leu Lys Val Lys Arg Leu Arg  
225 1 5 10  
227 <210> SEQ ID NO: 8  
228 <211> LENGTH: 12  
229 <212> TYPE: PRT  
230 <213> ORGANISM: Artificial Sequence  
232 <220> FEATURE:  
233 <223> OTHER INFORMATION: Peptide that binds a hyaluronan  
235 <221> NAME/KEY: HELIX  
236 <222> LOCATION: (1)...(5)  
237 <223> OTHER INFORMATION: Alpha-helix  
239 <400> SEQUENCE: 8  
240 Met Met Thr Val Leu Lys Val Lys Val Lys Arg Lys  
241 1 5 10  
243 <210> SEQ ID NO: 9  
244 <211> LENGTH: 10  
245 <212> TYPE: PRT  
246 <213> ORGANISM: Artificial Sequence  
248 <220> FEATURE:  
249 <223> OTHER INFORMATION: Peptide that binds a hyaluronan  
251 <221> NAME/KEY: HELIX  
252 <222> LOCATION: (1)...(5)  
253 <223> OTHER INFORMATION: Alpha-helix  
255 <400> SEQUENCE: 9  
256 Met Met Thr Val Leu Lys Val Arg Lys Arg  
257 1 5 10  
259 <210> SEQ ID NO: 10  
260 <211> LENGTH: 9  
261 <212> TYPE: PRT  
262 <213> ORGANISM: Artificial Sequence

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/685,010

DATE: 06/07/2001  
TIME: 11:23:29

Input Set : A:\401c1.app.txt  
Output Set: C:\CRF3\06072001\I685010.raw

264 <220> FEATURE:  
 265 <223> OTHER INFORMATION: Peptide that binds a hyaluronan  
 267 <221> NAME/KEY: HELIX  
 268 <222> LOCATION: (1)...(5)  
 269 <223> OTHER INFORMATION: Alpha-helix  
 271 <400> SEQUENCE: 10  
 272 Met Met Thr Val Leu Lys Val Arg Lys  
 273 1 5  
 275 <210> SEQ ID NO: 11  
 276 <211> LENGTH: 13  
 277 <212> TYPE: PRT  
 278 <213> ORGANISM: Homo sapiens  
 280 <400> SEQUENCE: 11  
 281 Lys Leu Gln Ala Thr Gln Lys Pro Leu Thr Glu Ser Lys  
 282 1 5 10  
 284 <210> SEQ ID NO: 12  
 285 <211> LENGTH: 12  
 286 <212> TYPE: PRT  
 287 <213> ORGANISM: Homo sapiens  
 289 <400> SEQUENCE: 12  
 290 Val Ser Ile Glu Lys Glu Lys Ile Asp Glu Lys Ser  
 291 1 5 10  
 293 <210> SEQ ID NO: 13  
 294 <211> LENGTH: 6  
 295 <212> TYPE: PRT  
 296 <213> ORGANISM: Artificial Sequence  
 298 <220> FEATURE:  
 299 <223> OTHER INFORMATION: Peptide developed based upon the TAM domain  
 300 (Transient Activator of MAP kinases)  
 302 <221> NAME/KEY: VARIANT  
 303 <222> LOCATION: (3)...(3)  
 304 <223> OTHER INFORMATION: Xaa = Any amino acid  
 306 <400> SEQUENCE: 13  
 307 Val Ser Xaa Lys Glu Lys  
 308 1 5  
 310 <210> SEQ ID NO: 14  
 311 <211> LENGTH: 23  
 312 <212> TYPE: PRT  
 313 <213> ORGANISM: Mus musculus  
 315 <400> SEQUENCE: 14  
 316 Lys Leu Gln Ala Thr Gln Lys Asp Leu Thr Glu Ser Lys Gly Lys Ile  
 317 1 5 10 15  
 318 Val Gln Leu Glu Gly Lys Leu  
 319 20  
 321 <210> SEQ ID NO: 15  
 322 <211> LENGTH: 14  
 323 <212> TYPE: PRT  
 324 <213> ORGANISM: Mus musculus  
 326 <400> SEQUENCE: 15

*FYT*  
**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

**VERIFICATION SUMMARY**  
PATENT APPLICATION: US/09/685,010

DATE: 06/07/2001  
TIME: 11:23:30

Input Set : A:\401c1.app.txt  
Output Set: C:\CRF3\06072001\I685010.raw

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:42 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:84 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:192 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:307 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13  
L:480 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28  
L:514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29

OIPE

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/09/637,054**DATE: 06/07/2001  
TIME: 11:29:04Input Set : A:\PTO.txt  
Output Set: C:\CRF3\06072001\I637054.raw**ENTERED**

3 <110> APPLICANT: Maroney, Anna  
4       Walton, Kevin M.  
5       Dionne, Craig A.  
6       Neff, Nicola  
7       Knight, Jr., Ernest  
8       Glicksman, Marcie A.  
10 <120> TITLE OF INVENTION: Methods For Modulating Multiple Lineage Kinase Proteins And Screening  
11       Compounds Which Modulate Lineage Kinase Proteins  
13 <130> FILE REFERENCE: CEPH-1235  
15 <140> CURRENT APPLICATION NUMBER: 09/637,054  
16 <141> CURRENT FILING DATE: 2000-08-11  
18 <160> NUMBER OF SEQ ID NOS: 18  
20 <170> SOFTWARE: PatentIn version 3.0  
22 <210> SEQ ID NO: 1  
23 <211> LENGTH: 17  
24 <212> TYPE: PRT  
25 <213> ORGANISM: Homo sapiens  
27 <400> SEQUENCE: 1  
29 Cys Gly Gly Ala Thr Cys Cys Ala Cys Met Gly Ile Gly Ala Tyr Tyr  
30 1                   5                   10                   15  
32 Thr  
35 <210> SEQ ID NO: 2  
36 <211> LENGTH: 23  
37 <212> TYPE: PRT  
38 <213> ORGANISM: Homo sapiens  
40 <400> SEQUENCE: 2  
42 Gly Gly Ala Ala Thr Thr Cys Cys Ala Trp Ala Gly Gly Ala Cys Cys  
43 1                   5                   10                   15  
45 Ala Ser Ala Cys Arg Thr Cys  
46                   20  
48 <210> SEQ ID NO: 3  
49 <211> LENGTH: 33  
50 <212> TYPE: PRT  
51 <213> ORGANISM: Homo sapiens  
53 <400> SEQUENCE: 3  
55 Cys Gly Gly Ala Thr Cys Cys Arg Thr Ile Cys Ala Tyr Met Gly Ile  
56 1                   5                   10                   15  
58 Gly Ala Tyr Tyr Thr Ile Gly Cys Ile Gly Cys Ile Met Gly Ile Ala  
59                   20                   25                   30  
61 Ala  
64 <210> SEQ ID NO: 4  
65 <211> LENGTH: 30  
66 <212> TYPE: PRT  
67 <213> ORGANISM: Homo sapiens  
69 <400> SEQUENCE: 4  
71 Gly Gly Ala Ala Thr Thr Ile Ala Tyr Ile Gly Gly Ala Trp Ala Ile  
72 1                   5                   10                   15

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/09/637,054**

**DATE: 06/07/2001**  
**TIME: 11:29:04**

**Input Set : A:\PTO.txt**  
**Output Set: C:\CRF3\06072001\I637054.raw**

145 <213> ORGANISM: Homo sapiens		
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151 <210> SEQ ID NO: 13		
152 <211> LENGTH: 8		
153 <212> TYPE: PRT		
154 <213> ORGANISM: Homo sapiens		
156 <400> SEQUENCE: 13		
158 Asp Tyr Lys Asp Asp Asp Asp Lys		
159 1 5		
161 <210> SEQ ID NO: 14		
162 <211> LENGTH: 69		
163 <212> TYPE: DNA		
164 <213> ORGANISM: Homo sapiens		
166 <400> SEQUENCE: 14		
167 ataaagcttc cagaggccat ggactacaag gacgacgatg acaaggcctg cctccatgaa	60	
169 acccgaaca	69	
172 <210> SEQ ID NO: 15		
173 <211> LENGTH: 18		
174 <212> TYPE: DNA		
175 <213> ORGANISM: Homo sapiens		
177 <400> SEQUENCE: 15		
178 gacagggcgg ccggctct	18	
181 <210> SEQ ID NO: 16		
182 <211> LENGTH: 583		
183 <212> TYPE: DNA		
184 <213> ORGANISM: Homo sapiens		
186 <400> SEQUENCE: 16		
187 gaattcggca cgagaggact cgcaggtgtc cggcgacgag ggctgggtga cccggcagct	60	
189 gaaccagcgg gtgggcatact tccccagcaa ctacgtgacc cccgcgcagcg ccttctccag	120	
191 cccgtgcag cccggcggcg aggaccccag ttgctaccgg cccattcaagt tgttagaaat	180	
193 tgattttgcg gagctcacct tggaaagagat tattggcatc gggggcttg ggaaggctta	240	
195 tcgtgctttc tggatagggg atgaggttgc tgtgaaagca gctcgccacg accctgatga	300	
197 ggacatcagc cagaccatag agaatgttgc ccaagaggcc aagctctcg ccatgctgaa	360	
199 gcaccccaac atcattgccc taagaggggt atgtctgaag gagcccaacc tctgcttgg	420	
201 catggagttt gctcggtggag gaccttggaa tagagtgtta tctggaaaa ggattcccc	480	
203 agacatcctg gtgaattggg ctgtgcagat tgccagaggg atgaactact tacatgatga	540	
205 ggcaatttgtt cccatcatcc accgcgaccc taagtccagc aac	583	
208 <210> SEQ ID NO: 17		
209 <211> LENGTH: 194		
210 <212> TYPE: PRT		
211 <213> ORGANISM: Homo sapiens		
213 <400> SEQUENCE: 17		
215 Asn Ser Ala Arg Glu Asp Ser Gln Val Ser Gly Asp Glu Gly Trp Trp		
216 1 5 10 15		
218 Thr Gly Gln Leu Asn Gln Arg Val Gly Ile Phe Pro Ser Asn Tyr Val		
219 20 25 30		
221 Thr Pro Arg Ser Ala Phe Ser Ser Arg Cys Gln Pro Gly Gly Glu Asp		
222 35 40 45		

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/637,054

DATE: 06/07/2001  
TIME: 11:29:04

Input Set : A:\PTO.txt  
Output Set: C:\CRF3\06072001\I637054.raw

224 Pro Ser Cys Tyr Pro Pro Ile Gln Leu Leu Glu Ile Asp Phe Ala Glu  
225 50 55 60  
227 Leu Thr Leu Glu Glu Ile Ile Gly Ile Gly Gly Phe Gly Lys Val Tyr  
228 65 70 75 80  
230 Arg Ala Phe Trp Ile Gly Asp Glu Val Ala Val Lys Ala Ala Arg His  
231 85 90 95  
233 Asp Pro Asp Glu Asp Ile Ser Gln Thr Ile Glu Asn Val Arg Gln Glu  
234 100 105 110  
236 Ala Lys Leu Phe Ala Met Leu Lys His Pro Asn Ile Ile Ala Leu Arg  
237 115 120 125  
239 Gly Val Cys Leu Lys Glu Pro Asn Leu Cys Leu Val Met Glu Phe Ala  
240 130 135 140  
242 Arg Gly Gly Pro Leu Asn Arg Val Leu Ser Gly Lys Arg Ile Pro Pro  
243 145 150 155 160  
245 Asp Ile Leu Val Asn Trp Ala Val Gln Ile Ala Arg Gly Met Asn Tyr  
246 165 170 175  
248 Leu His Asp Glu Ala Ile Val Pro Ile Ile His Arg Asp Leu Lys Ser  
249 180 185 190  
251 Ser Asn  
254 <210> SEQ ID NO: 18  
255 <211> LENGTH: 10  
256 <212> TYPE: PRT  
257 <213> ORGANISM: Homo sapiens  
259 <400> SEQUENCE: 18  
261 Asn Asp Tyr Lys Asp Asp Asp Asp Lys Cys  
262 1 5 10

**VERIFICATION SUMMARY**  
PATENT APPLICATION: US/09/637,054

DATE: 06/07/2001  
TIME: 11:29:05

Input Set : A:\PTO.txt  
Output Set: C:\CRF3\06072001\I637054.raw



Creation date: 11-17-2003  
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Legal Date: 11-07-2001

No.	Doccode	Number of pages
1	PA..	3

Total number of pages: 3

Remarks:

Order of re-scan issued on .....